

O I P E

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SEQUENCE LISTING

1635

PATENT & TRADEMARK OFFICE

Sheet 6
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JONES, Stacey A.
WILLSON, Timothy M.

<120> AN ORPHAN NUCLEAR RECEPTOR

<130> 510-125

<140> 09/276,935

<141> 1999-03-26

<150> 60/079,593

<151> 1998-03-27

<160> 14

<170> PatentIn Ver. 2.0

<210> 1

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: DNA genome

<400> 1

ctgctgcgca tccaggacat

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<210> 2

<211> 45

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: DNA genome

<400> 2

gggtgtgggg aatccaccac catggaggtg agacccaaag aaagc

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<210> 3

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: DNA genome

<400> 3

gggtgtgggg gatcctcagc tacctgttat gccc

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<210> 4

<211> 31

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: DNA genome

<400> 4
gatcagacag ttcatgaagt tcatcttagat c

<210> 5
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: DNA genome

<400> 5
gatcaatatg aactcaaagg aggtcagtg

<210> 6
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<220>
<223> Description of Artificial Sequence: DNA genome

<400> 6
gatcaatatg aactcaaagg aggtcagtg

<210> 7
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: DNA genome

<400> 7
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<210> 8
<211> 29
<212> DNA
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<220>
<223> Description of Artificial Sequence: DNA genome

<400> 8
gatcaataac aactcaaagg aggtcagtg

<210> 9
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<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: DNA genome

<400> 9
gatgcagaca gttcatgaag ttcatctaga tc

<210> 10
<211> 11
<212> PRT

31

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32

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Protein

<400> 10

Met Lys Lys Gly His His His His His His Gly
1 5 10

<210> 11

<211> 316

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Protein

<400> 11

Met Lys Lys Gly His His His His His His Gly Ser Glu Arg Thr Gly
1 5 10 15

Thr Gln Pro Leu Gly Val Gln Gly Leu Thr Glu Glu Gln Arg Met Met
20 25 30

Ile Arg Glu Leu Met Asp Ala Gln Met Lys Thr Phe Asp Thr Thr Phe
35 40 45

Ser His Phe Lys Asn Phe Arg Leu Pro Gly Val Leu Ser Ser Gly Cys
50 55 60

Glu Leu Pro Glu Ser Leu Gln Ala Pro Ser Arg Glu Glu Ala Ala Lys
65 70 75 80

Trp Ser Gln Val Arg Lys Asp Leu Cys Ser Leu Lys Val Ser Leu Gln
85 90 95

Leu Arg Gly Glu Asp Gly Ser Val Trp Asn Tyr Lys Pro Pro Ala Asp
100 105 110

Ser Gly Gly Lys Glu Ile Phe Ser Leu Leu Pro His Met Ala Asp Met
115 120 125

Ser Thr Tyr Met Phe Lys Gly Ile Ile Ser Phe Ala Lys Val Ile Ser
130 135 140

Tyr Phe Arg Asp Leu Pro Ile Glu Asp Gln Ile Ser Leu Leu Lys Gly
145 150 155 160

Ala Ala Phe Glu Leu Cys Gln Leu Arg Phe Asn Thr Val Phe Asn Ala
165 170 175

Glu Thr Gly Thr Trp Glu Cys Gly Arg Leu Ser Tyr Cys Leu Glu Asp
180 185 190

Thr Ala Gly Gly Phe Gln Gln Leu Leu Leu Glu Pro Met Leu Lys Phe
195 200 205

His Tyr Met Leu Lys Lys Leu Gln Leu His Glu Glu Tyr Val Leu
210 215 220

Met Gln Ala Ile Ser Leu Phe Ser Pro Asp Arg Pro Gly Val Leu Gln
225 230 235 240

His Arg Val Val Asp Gln Leu Gln Glu Gln Phe Ala Ile Thr Leu Lys
245 250 255

Ser Tyr Ile Glu Cys Asn Arg Pro Gln Pro Ala His Arg Phe Leu Phe
260 265 270

Leu Lys Ile Met Ala Met Leu Thr Glu Leu Arg Ser Ile Asn Ala Gln
275 280 285

His Thr Gln Arg Leu Leu Arg Ile Gln Asp Ile His Pro Phe Ala Thr
290 295 300

Pro Leu Met Gln Glu Leu Phe Gly Ile Thr Gly Ser
305 310 315

<210> 12

<211> 242

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Protein

<400> 12

Met Lys Lys Gly Ser Ala Asn Glu Asp Met Pro Val Glu Arg Ile Leu
1 5 10 15

Glu Ala Glu Leu Ala Val Glu Pro Lys Thr Glu Thr Tyr Val Glu Ala
20 25 30

Asn Met Gly Leu Asn Pro Ser Ser Pro Asn Asp Pro Val Thr Asn Ile
35 40 45

Cys Gln Ala Ala Asp Lys Gln Leu Phe Thr Leu Val Glu Trp Ala Lys
50 55 60

Arg Ile Pro His Phe Ser Glu Leu Pro Leu Asp Asp Gln Val Ile Leu
65 70 75 80

Leu Arg Ala Gly Trp Asn Glu Leu Leu Ile Ala Ser Phe Ser His Arg
85 90 95

Ser Ile Ala Val Lys Asp Gly Ile Leu Leu Ala Thr Gly Leu His Val
100 105 110

His Arg Asn Ser Ala His Ser Ala Gly Val Gly Ala Ile Phe Asp Arg
115 120 125

Val Leu Thr Glu Leu Val Ser Lys Met Arg Asp Met Gln Met Asp Lys
130 135 140

Thr Glu Leu Gly Cys Leu Arg Ala Ile Val Leu Phe Asn Pro Asp Ser
145 150 155 160

Lys Gly Leu Ser Asn Pro Ala Glu Val Glu Ala Leu Arg Glu Lys Val
165 170 175

Tyr Ala Ser Leu Glu Ala Tyr Cys Lys His Lys Tyr Pro Glu Gln Pro
180 185 190

Gly Arg Phe Ala Lys Leu Leu Leu Arg Leu Pro Ala Leu Arg Ser Ile
195 200 205

Gly Leu Lys Cys Leu Glu His Leu Phe Phe Phe Lys Leu Ile Gly Asp
210 215 220

Thr Pro Ile Asp Thr Phe Leu Met Glu Met Leu Glu Ala Pro His Gln
225 230 235 240

Met Thr

<210> 13

<211> 2146

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: DNA genome

<400> 13

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aagtgttac agtgagaaaa gcaagagaat aacttataac tcctgtctg aacaaggcag 180
cggttccttg gttaagctac tccttgcattc atccatttgc caaggattttt ccaaagtggac 240
cccaggggag aagtccggagc aaagaactta ccaccacgca gtccaaaggagg cccagaagca 300
aacctggagg tgagacccaa agaaagctgg aaccatgtc actttgtaca ctgtgaggac 360
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atctgcctgt tatgtgggaa caaggccact ggttataact tcaatgtcat gacatgtgaa 480
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cggaaggggcg cctgcgagat cacccggaaag accccggcgc acgtggccagc ctgcggcc 600
cgcaagtgcc tggagagccg catagaagaag gagatgtatc tgccgacga ggccgtggag 660
gagaggccgg cttgtatcaa gcggaaagaa agtgcacgga caggactca gccactggga 720
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ccatctgggg tctatgcctca cataccacg tttgttgcgt tcctgttgc tttcattgtc 2040
acctctaata gtccctgtctc ccacttccca ctgcgttcccc tcctcttccg agctgtttt 2100
tgggtccag gcctgtactc atccggcaggat gcatgagtat ctgttgc 2146

<210> 14
<211> 414
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Protein

<400> 14
Leu Glu Val Arg Pro Lys Glu Ser Trp Asn His Ala Asp Phe Val His
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Cys Glu Asp Thr Glu Ser Val Pro Gly Lys Pro Ser Val Asn Ala Asp
20 25 30

Glu Glu Val Gly Gly Pro Gln Ile Cys Arg Val Cys Gly Asp Lys Ala
35 40 45

Thr Gly Tyr His Phe Asn Val Met Thr Cys Glu Gly Cys Lys Gly Phe
50 55 60

Phe Arg Arg Ala Met Lys Arg Asn Ala Arg Leu Arg Cys Pro Phe Arg
65 70 75 80

Lys Gly Ala Cys Glu Ile Thr Arg Lys Thr Arg Arg Gln Cys Gln Ala
85 90 95

Cys Arg Leu Arg Lys Cys Leu Glu Ser Gly Met Lys Lys Glu Met Ile
100 105 110

Met Ser Asp Glu Ala Val Glu Glu Arg Arg Ala Leu Ile Lys Arg Lys
115 120 125

Lys Ser Glu Arg Thr Gly Thr Gln Pro Leu Gly Val Gln Gly Leu Thr
130 135 140

Glu Glu Gln Arg Met Met Ile Arg Glu Leu Met Asp Ala Gln Met Lys
145 150 155 160

Thr Phe Asp Thr Phe Ser His Phe Lys Asn Phe Arg Leu Pro Gly
165 170 175

Val Leu Ser Ser Gly Cys Glu Leu Pro Glu Ser Leu Gln Ala Pro Ser
180 185 190

Arg Glu Glu Ala Ala Lys Trp Ser Gln Val Arg Lys Asp Leu Cys Ser
195 200 205

Leu Lys Val Ser Leu Gln Leu Arg Gly Glu Asp Gly Ser Val Trp Asn
210 215 220

Tyr Lys Pro Pro Ala Asp Ser Gly Gly Lys Glu Ile Phe Ser Leu Leu
225 230 235 240

Pro His Met Ala Asp Met Ser Thr Tyr Met Phe Lys Gly Ile Ile Ser
245 250 255

Phe Ala Lys Val Ile Ser Tyr Phe Arg Asp Leu Pro Ile Glu Asp Gln
260 265 270

Ile Ser Leu Leu Lys Gly Ala Ala Phe Glu Leu Cys Gln Leu Arg Phe
275 280 285

Asn Thr Val Phe Asn Ala Glu Thr Gly Thr Trp Glu Cys Gly Arg Leu
290 295 300

Ser Tyr Cys Leu Glu Asp Thr Ala Gly Gly Phe Gln Gln Leu Leu Leu
305 310 315 320

Glu Pro Met Leu Lys Phe His Tyr Met Leu Lys Lys Leu Gln Leu His
325 330 335

Glu Glu Glu Tyr Val Leu Met Gln Ala Ile Ser Leu Phe Ser Pro Asp
340 345 350

Arg Pro Gly Val Leu Gln His Arg Val Val Asp Gln Leu Gln Glu Gln
355 360 365

Phe Ala Ile Thr Leu Lys Ser Tyr Ile Glu Cys Asn Arg Pro Gln Pro
370 375 380

Ala His Arg Phe Leu Phe Leu Lys Ile Met Ala Met Leu Thr Glu Phe
385 390 395 400

Ala Thr Pro Leu Met Gln Glu Leu Phe Gly Ile Thr Gly Ser
405 410